1: 3:		Gly	Pro	Gly	Arg	Gly	Pro	Val	Gly	Arg	Arg	Arg	Tyr	<u>Ma</u>	Arg	Lys	16
):						• • •	 Phe			 Lys			 His	Val Pro	 Lys		
3 : A :	Gln	Leu	Val	Pro	Leu	Leu	Tyr	Lys	G1n	Phe	Val	Pro	<u>617</u>	Val	Pro	Glu	32
C: D:			 Thr		• • • •	 Ala					 Ile		Ser Asn	Met	 Ala	• • • •	
E: A:	Arg	Thr	Leu	Gly	Ala	Ser	Gly	Pro	Ala	Glu	G1y	Arg	Val	Ale	Arg	Gly	48
B: C: D: E:		•••	• • • •				• • •	 Arg	 Tyr			 Lys	 Ile	Thr Ser		 Asn	
A:		G1 u	Arg	Phe	Arg	Asp	Leu	Val	Pro	Asn	Tyr	Asn	Pro	Asp	Ile		64
				• • •													
A:	Phe	Lys	. Asp	Glu	G1·u	Asn	Ser	G1 y	Ala	Asp	Arg	Leu	Met	Thr	Glu	Arg	80
C: D:							 Thr						• • •		 Gln		٠
A:	Cys	Lys	s Glu	ı Arg	Val	Asn	Ala	Leu	Ala	Ile	Ala	. Val	Met	Asn	Met	Trp	96
C: D:	• • • •		Asp	 Lys	 Leu	٠٠٠					 Ser	• • • •		• • •	 Gln	• • • •	
A:	Pro	o Gly	y Val	l Arg	, Lei	ı Arg	g Val	Thr	Glu	Gly	Trp) Asp	o Glu	ı Asp	o Gly	His	112
C: D:				Lys			• • •			• • •	• • •	• • •				• • •	
				. Lys n Ast												Thr	128
B: C: D:		 . Se		 u Glu			• • • •	• • •	 		• • •	• • • •	 	 l	 	• • • •	
Ē :		. Se	r Gl	u Gli	ı						• • •		. Va	1	• • • •		

																Ala	144
E :														• • •	• • •	• • •	
	•																
A:	Val	Glu	Ala	G1 y	Phe	Asp	Trp	Val	Tyr	Tyr	Glu	Ser	Arg	Asn	His	Ile	160
B :					•.••						Gly						
E :													Lys	Ala	• • •	Val	
_		Val	Ser	Val	Lys	Ala	Asp	Asn	Ser	Leu	Ala	Val	Arg	Ala	G1y	Gly	176
B:							,					•					
Ľ.	• • •	Cys				001	olu	1113		11 T C	• • •	,, <u>, , , , , , , , , , , , , , , , , ,</u>	2,0				

REFERENCE FIGURE 1. Sequence comparison of hedgehog proteins by this invention and others.

A, human mature Desert hedgehog protein by this invtion (SEQ ID NO:1=residues Cys1-Gly176 of amino acid sequence alinged with SEQ ID NO:7); B, a part of human Desert hedgehog protein by Drummond; C, mouse Desert hedgehog protein by Ingham et al. (US PAT NO.5,884,079, SEQ ID NO:9, Cys23-Gly198); D, human Sonic hedgehog protein by Ingham et al. (US'079, SEQ ID NO:13, Cys24-Gly197); E, a part of human Indian hedgehog protein by Ingham et al. (US'079, SEQ ID NO:14, Arg1-Gly103).

Triplet dot means an amino acid residue identical with that of the sequence A. At the position with parentheses, a gap is introduced. Unidentified portions are shown as thin lines.